SEQUENCE LISTING

<110> Croce, Carlo
Brenner, Charles
Pekarski,Yuri

<120> CRYSTAL STRUCTURE OF WORM NitFhit
REVEALS THAT A Nit TETRAMER BINDS TWO Fhit DIMERS

<130> CRO01.NP007

<150> 60/204,713
<151> 2000-05-16

<160> 11

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 276
<212> PRT
<213> Homo sapien

<400> 1

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 1               5                  10                  15
Lys Ser Asp Asn Val Thr Arg Ala Cys Ser Phe Ile Arg Glu Ala Ala
            20                  25                  30
Thr Gln Gly Ala Lys Ile Val Ser Leu Pro Glu Cys Phe Asn Ser Pro
        35                  40                  45
Tyr Gly Ala Lys Tyr Phe Pro Glu Tyr Ala Glu Lys Ile Pro Gly Glu
    50                  55                  60
Ser Thr Gln Lys Leu Ser Glu Val Ala Lys Glu Cys Ser Ile Tyr Leu
65                  70                  75                  80
Ile Gly Gly Ser Ile Pro Glu Glu Asp Ala Gly Lys Leu Tyr Asn Thr
                85                  90                  95
Cys Ala Val Phe Gly Pro Asp Gly Thr Leu Leu Ala Lys Tyr Arg Lys
            100                 105                 110
Ile His Leu Phe Asp Ile Asp Val Pro Gly Lys Ile Thr Phe Gln Glu
        115                 120                 125
Ser Lys Thr Leu Ser Pro Gly Asp Ser Phe Ser Thr Phe Asp Thr Pro
    130                 135                 140
Tyr Cys Arg Val Gly Leu Gly Ile Cys Tyr Asp Met Arg Phe Ala Glu
145                 150                 155                 160
Leu Ala Gln Ile Tyr Ala Gln Arg Gly Cys Gln Leu Leu Val Tyr Pro
                165                 170                 175
Gly Ala Phe Asn Leu Thr Thr Gly Pro Ala His Trp Glu Leu Leu Gln
            180                 185                 190
Arg Ser Arg Ala Val Asp Asn Gln Val Tyr Val Ala Thr Ala Ser Pro
        195                 200                 205
Ala Arg Asp Asp Lys Ala Ser Tyr Val Ala Trp Gly His Ser Thr Val
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    210                 215                 220
Val Asn Pro Trp Gly Glu Val Leu Ala Lys Ala Gly Thr Glu Glu Ala
225                 230                 235                 240
Ile Val Tyr Ser Asp Ile Asp Leu Lys Lys Leu Ala Glu Ile Arg Gln
                245                 250                 255
Gln Ile Pro Val Phe Arg Gln Lys Arg Ser Asp Leu Tyr Ala Val Glu
            260                 265                 270
Met Lys Lys Pro
        275
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<210> 2
<211> 276
<212> PRT
<213> mouse

<400> 2

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Met Ser Thr Phe Arg Leu Ala Leu Ile Gln Leu Gln Val Ser Ser Ile
 1                5                  10                  15
Lys Ser Asp Asn Leu Thr Arg Ala Cys Ser Leu Val Arg Glu Ala Ala
            20                  25                  30
Lys Gln Gly Ala Asn Ile Val Ser Leu Pro Glu Cys Phe Asn Ser Pro
        35                  40                  45
Tyr Gly Thr Thr Tyr Phe Pro Asp Tyr Ala Glu Lys Ile Pro Gly Glu
    50                  55                  60
Ser Thr Gln Lys Leu Ser Glu Val Ala Lys Glu Ser Ser Ile Tyr Leu
65                  70                  75                  80
Ile Gly Gly Ser Ile Pro Glu Glu Asp Ala Gly Lys Leu Tyr Asn Thr
                85                  90                  95
Cys Ser Val Phe Gly Pro Asp Gly Ser Leu Leu Val Lys His Arg Lys
            100                 105                 110
Ile His Leu Phe Asp Ile Asp Val Pro Gly Lys Ile Thr Phe Gln Glu
        115                 120                 125
Ser Lys Thr Leu Ser Pro Gly Asp Ser Phe Ser Thr Phe Asp Thr Pro
    130                 135                 140
Tyr Cys Lys Val Gly Leu Gly Ile Cys Tyr Asp Met Arg Phe Ala Glu
145                 150                 155                 160
Leu Ala Gln Ile Tyr Ala Gln Arg Gly Cys Gln Leu Leu Val Tyr Pro
                165                 170                 175
Gly Ala Phe Asn Leu Thr Thr Gly Pro Ala His Trp Glu Leu Leu Gln
            180                 185                 190
Arg Ala Arg Ala Val Asp Asn Gln Val Tyr Val Ala Thr Ala Ser Pro
        195                 200                 205
Ala Arg Asp Asp Lys Ala Ser Tyr Val Ala Trp Gly His Ser Thr Val
    210                 215                 220
Val Asp Pro Trp Gly Gln Val Leu Thr Lys Ala Gly Thr Glu Glu Thr
225                 230                 235                 240
Ile Leu Tyr Ser Asp Ile Asp Leu Lys Lys Leu Ala Glu Ile Arg Gln
                245                 250                 255
Gln Ile Pro Ile Leu Lys Gln Lys Arg Ala Asp Leu Tyr Thr Val Glu
            260                 265                 270
Ser Lys Lys Pro
        275
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<210> 3
<211> 288

<212> PRT
<213> X. laevis

<400> 3
Met Ala Gly Ala His Lys Pro Leu Ile Ala Val Cys Gln Met Thr Ser
 1   5   10   15
Thr Ser Asp Lys Glu Lys Asn Phe Ala Thr Cys Ser Arg Leu Ile Arg
            20   25   30
Glu Ala Ala Gly Arg Arg Ala Cys Met Val Phe Leu Pro Glu Ala Phe
        35   40   45
Asp Tyr Ile Gly Gly Ser Ile Glu Glu Thr Leu Ser Leu Ala Glu Ser
    50   55   60
Leu His Gly Asp Thr Ile Gln Arg Tyr Thr Gln Leu Ala Arg Glu Cys
65   70   75   80
Gly Leu Trp Leu Ser Leu Gly Gly Phe His Glu Lys Gly Pro Asn Trp
                85   90   95
Asp Thr Asp Gln Arg Ile Ser Asn Ser His Val Val Val Asp Asn Thr
            100   105   110
Gly His Ile Val Ser Val Tyr Arg Lys Ala His Leu Phe Asp Val Asp
        115   120   125
Leu Gln Asn Gly Val Ser Leu Arg Glu Ser Ser Ser Thr Leu Pro Gly
    130   135   140
Ala Glu Leu Ile Arg Pro Ile Thr Ser Pro Ala Gly Lys Ile Gly Leu
145   150   155   160
Gly Val Cys Tyr Asp Leu Arg Phe Pro Glu Phe Ser Leu Ala Leu Ala
                165   170   175
Gln Gln Gly Ala Glu Leu Leu Thr Tyr Pro Ser Ala Phe Thr Leu Thr
            180   185   190
Thr Gly Leu Ala His Trp Glu Val Leu Leu Arg Ala Arg Ala Ile Glu
        195   200   205
Thr Gln Cys Tyr Val Val Ala Ala Ala Gln Thr Asp Arg His Asn Glu
    210   215   220
Lys Arg Thr Ser Tyr Gly His Ala Met Val Val Asp Pro Trp Gly Leu
225   230   235   240
Val Ile Gly Gln Cys Gln Glu Gly Thr Gly Ile Cys Tyr Ala Glu Ile
                245   250   255
Asp Ile Pro Tyr Met Glu Arg Val Arg Arg Asp Met Pro Val Trp Arg
            260   265   270
His Arg Arg Thr Asp Leu Tyr Gly Lys Ile Ser Phe Asn Lys Pro Asp
        275   280   285

<210> 4
<211> 307
<212> PRT
<213> S. cerevisiae

<400> 4
Met Thr Ser Lys Leu Lys Arg Val Ala Val Ala Gln Leu Cys Ser Ser
 1   5   10   15
Ala Asp Leu Thr Lys Asn Leu Lys Val Val Lys Glu Leu Ile Ser Glu
            20   25   30
Ala Ile Gln Lys Lys Ala Asp Val Val Phe Leu Pro Glu Ala Ser Asp
        35   40   45
Tyr Leu Ser Gln Asn Pro Leu His Ser Arg Tyr Leu Ala Gln Lys Ser
    50   55   60

```
Pro Lys Phe Ile Arg Gln Leu Gln Ser Ser Ile Thr Asp Leu Val Arg
65                  70                  75                  80
Asp Asn Ser Arg Asn Ile Asp Val Ser Ile Gly Val His Leu Pro Pro
                85                  90                  95
Ser Glu Gln Asp Leu Leu Glu Gly Asn Asp Arg Val Arg Asn Val Leu
            100                 105                 110
Leu Tyr Ile Asp His Glu Gly Lys Ile Leu Gln Glu Tyr Gln Lys Leu
        115                 120                 125
His Leu Phe Asp Val Asp Val Pro Asn Gly Pro Ile Leu Lys Glu Ser
    130                 135                 140
Lys Ser Val Gln Pro Gly Lys Ala Ile Pro Asp Ile Ile Glu Ser Pro
145                 150                 155                 160
Leu Gly Lys Leu Gly Ser Ala Ile Cys Tyr Asp Ile Arg Phe Pro Glu
                165                 170                 175
Phe Ser Leu Lys Leu Arg Ser Met Gly Ala Glu Ile Leu Cys Phe Pro
            180                 185                 190
Ser Ala Phe Thr Ile Lys Thr Gly Glu Ala His Trp Glu Leu Leu Gly
        195                 200                 205
Arg Ala Arg Ala Val Asp Thr Gln Cys Tyr Val Leu Met Pro Gly Gln
    210                 215                 220
Val Gly Met His Asp Leu Ser Asp Pro Glu Trp Glu Lys Gln Ser His
225                 230                 235                 240
Met Ser Ala Leu Glu Lys Ser Ser Arg Arg Glu Ser Trp Gly His Ser
                245                 250                 255
Met Val Ile Asp Pro Trp Gly Lys Ile Ile Ala His Ala Asp Pro Ser
            260                 265                 270
Thr Val Gly Pro Gln Leu Ile Leu Ala Asp Leu Asp Arg Glu Leu Leu
        275                 280                 285
Gln Glu Ile Arg Asn Lys Met Pro Leu Trp Asn Gln Arg Arg Asp Asp
    290                 295                 300
Leu Phe His
305
```

<210> 5
<211> 291
<212> PRT
<213> S. cerevisiae

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<400> 5
Met Ser Ala Ser Lys Ile Leu Ser Gln Lys Ile Lys Val Ala Leu Val
 1               5                  10                  15
Gln Leu Ser Gly Ser Ser Pro Asp Lys Met Ala Asn Leu Gln Arg Ala
            20                  25                  30
Ala Thr Phe Ile Glu Arg Ala Met Lys Glu Gln Pro Asp Thr Lys Leu
        35                  40                  45
Val Val Leu Pro Glu Cys Phe Asn Ser Pro Tyr Ser Thr Asp Gln Phe
    50                  55                  60
Arg Lys Tyr Ser Glu Val Ile Asn Pro Lys Glu Pro Ser Thr Ser Val
65                  70                  75                  80
Gln Phe Leu Ser Asn Leu Ala Asn Lys Phe Lys Ile Ile Leu Val Gly
                85                  90                  95
Gly Thr Ile Pro Glu Leu Asp Pro Lys Thr Asp Lys Ile Tyr Asn Thr
            100                 105                 110
Ser Ile Ile Phe Asn Glu Asp Gly Lys Leu Ile Asp Lys His Arg Lys
        115                 120                 125
```

Val His Leu Phe Asp Val Asp Ile Pro Asn Gly Ile Ser Phe His Glu
130 135 140
Ser Glu Thr Leu Ser Pro Gly Glu Lys Ser Thr Thr Ile Asp Thr Lys
145 150 155 160
Tyr Gly Lys Phe Gly Val Gly Ile Cys Tyr Asp Met Arg Phe Pro Glu
165 170 175
Leu Ala Met Leu Ser Ala Arg Lys Gly Ala Phe Ala Met Ile Tyr Pro
180 185 190
Ser Ala Phe Asn Thr Val Thr Gly Pro Leu His Trp His Leu Leu Ala
195 200 205
Arg Ser Arg Ala Val Asp Asn Gln Val Tyr Val Met Leu Cys Ser Pro
210 215 220
Ala Arg Asn Leu Gln Ser Ser Tyr His Ala Tyr Gly His Ser Ile Val
225 230 235 240
Val Asp Pro Arg Gly Lys Ile Val Ala Glu Ala Gly Glu Gly Glu Glu
245 250 255
Ile Ile Tyr Ala Glu Leu Asp Pro Glu Val Ile Glu Ser Phe Arg Gln
260 265 270
Ala Val Pro Leu Thr Lys Gln Arg Arg Phe Asp Val Tyr Ser Asp Val
275 280 285
Asn Ala His
290

<210> 6
<211> 276
<212> PRT
<213> S. pombe

<400> 6
Met Thr Leu Ala Ala Val Ala Gln Leu Asn Ser Ser Gly Ser Ile Leu
1 5 10 15
Lys Asn Leu Ala Ile Cys Lys Glu Leu Ile Ser Gln Ala Ala Ala Lys
20 25 30
Gly Ala Lys Cys Ile Phe Phe Pro Glu Ala Ser Asp Phe Ile Ala His
35 40 45
Asn Ser Asp Glu Ala Ile Glu Leu Thr Asn His Pro Asp Cys Ser Lys
50 55 60
Phe Ile Arg Asp Val Arg Glu Ser Ala Thr Lys His Ser Ile Phe Val
65 70 75 80
Asn Ile Cys Val His Glu Pro Ser Lys Val Lys Asn Lys Leu Leu Asn
85 90 95
Ser Ser Leu Phe Ile Glu Pro Leu His Gly Glu Ile Ile Ser Arg Tyr
100 105 110
Ser Lys Ala His Leu Phe Asp Val Glu Ile Lys Asn Gly Pro Thr Leu
115 120 125
Lys Glu Ser Asn Thr Thr Leu Arg Gly Glu Ala Ile Leu Pro Pro Cys
130 135 140
Lys Thr Pro Leu Gly Lys Val Gly Ser Ala Ile Cys Phe Asp Ile Arg
145 150 155 160
Phe Pro Glu Gln Ala Ile Lys Leu Arg Asn Met Gly Ala His Ile Ile
165 170 175
Thr Tyr Pro Ser Ala Phe Thr Glu Lys Thr Gly Ala Ala His Trp Glu
180 185 190
Val Leu Leu Arg Ala Arg Ala Leu Asp Ser Gln Cys Tyr Val Ile Ala
195 200 205

Pro Ala Gln Gly Gly Lys His Asn Glu Lys Arg Ala Ser Tyr Gly His
210 215 220
Ser Met Ile Val Asp Pro Trp Gly Thr Val Ile Ala Gln Tyr Ser Asp
225 230 235 240
Ile Ser Ser Pro Asn Gly Leu Ile Phe Ala Asp Leu Asp Leu Asn Leu
245 250 255
Val Asp His Val Arg Thr Tyr Ile Pro Leu Leu Arg Arg Asn Asp Leu
260 265 270
Tyr Pro Thr Ile
275

<210> 7
<211> 322
<212> PRT
<213> S. pombe

<400> 7
Met Asn Ser Lys Phe Phe Gly Leu Val Gln Lys Gly Thr Arg Ser Phe
1 5 10 15
Phe Pro Ser Leu Asn Phe Cys Tyr Thr Arg Asn Ile Met Ser Val Ser
20 25 30
Ala Ser Ser Leu Val Pro Lys Asp Phe Arg Ala Phe Arg Ile Gly Leu
35 40 45
Val Gln Leu Ala Asn Thr Lys Asp Lys Ser Glu Asn Leu Gln Leu Ala
50 55 60
Arg Leu Lys Val Leu Glu Ala Ala Lys Asn Gly Ser Asn Val Ile Val
65 70 75 80
Leu Pro Glu Ile Phe Asn Ser Pro Tyr Gly Thr Gly Tyr Phe Asn Gln
85 90 95
Tyr Ala Glu Pro Ile Glu Glu Ser Ser Pro Ser Tyr Gln Ala Leu Ser
100 105 110
Ser Met Ala Lys Asp Thr Lys Thr Tyr Leu Phe Gly Gly Ser Ile Pro
115 120 125
Glu Arg Lys Asp Gly Lys Leu Tyr Asn Thr Ala Met Val Phe Asp Pro
130 135 140
Ser Gly Lys Leu Ile Ala Val His Arg Lys Ile His Leu Phe Asp Ile
145 150 155 160
Asp Ile Pro Gly Gly Val Ser Phe Arg Glu Ser Asp Ser Leu Ser Pro
165 170 175
Gly Asp Ala Met Thr Met Val Asp Thr Glu Tyr Gly Lys Phe Gly Leu
180 185 190
Gly Ile Cys Tyr Asp Ile Arg Phe Pro Glu Leu Ala Met Ile Ala Ala
195 200 205
Arg Asn Gly Cys Ser Val Met Ile Tyr Pro Gly Ala Phe Asn Leu Ser
210 215 220
Thr Gly Pro Leu His Trp Glu Leu Leu Ala Arg Ala Arg Ala Val Asp
225 230 235 240
Asn Glu Met Phe Val Ala Cys Cys Ala Pro Ala Arg Asp Met Asn Ala
245 250 255
Asp Tyr His Ser Trp Gly His Ser Thr Val Val Asp Pro Phe Gly Lys
260 265 270
Val Ile Ala Thr Thr Asp Glu Lys Pro Ser Ile Val Tyr Ala Asp Ile
275 280 285
Asp Pro Ser Val Met Ser Thr Ala Arg Asn Ser Val Pro Ile Tyr Thr
290 295 300

Gln Arg Arg Phe Asp Val Tyr Ser Glu Val Leu Pro Ala Leu Lys Lys
305 310 315 320
Glu Glu

<210> 8
<211> 1359
<212> DNA
<213> Homo sapien

<400> 8
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caaggagcca aaatagtttc tttgccggaa tgctttaatt ctccatatgg agcgaaatat 180
tttcctgaat atgcagagaa aattcctggt gaatccacac agaagctttc tgaagtagca 240
aaggaatgca gcatatatct cattggaggc tctatccctg aagaggatgc tgggaaatta 300
tataacacct gtgctgtgtt tgggcctgat ggaactttac tagcaaagta tagaaagatc 360
catctgtttg acattgatgt tcctggaaaa attacatttc aagaatctaa aacattgagt 420
ccgggtgata gtttctccac atttgatact ccttactgca gagtgggtct gggcatctgc 480
tacgacatgc ggtttgcaga gcttgcacaa atctacgcac agagaggctg ccagctgttg 540
gtatatccag gagcttttaa tctgaccact ggaccagccc attgggagtt acttcagcga 600
agccgggctg ttgataatca ggtgtatgtg gccacagcct ctcctgcccg ggatgacaaa 660
gcctcctatg ttgcctgggg acacagcacc gtggtgaacc cttgggggga ggttctagcc 720
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aaaaagccct aaagtttatg tttctaatgt gtcacagaat aggacgatat gattctacaa 900
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tggtgattga ttcacctaat ataaatatat ttgtgtcatg aacctcttaa aaagttgctg 1200
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ggtaccttgn aagttccatt tttgggttca tgagtagcta taggaacgca agggtgatac 1320
atctttgggt gttttgccag agaagttggg cagccccac 1359

<210> 9
<211> 1292
<212> DNA
<213> mouse

<400> 9
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agcctagtgc gggaggcagc aaagcaaggt gccaacatag tttctctgcc tgagtgcttc 180
aattctccat atggaacaac ctactttcct gactatgcag agaagattcc tggagagtcc 240
acacaaaagc tttctgaagt agcaaaggag agcagcatat atctcattgg aggctccatc 300
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gcccactggg agctgcttca gcgagcccgg gctgttgata atcaggtgta tgtggctaca 660
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gatccttggg ggcaggtcct aaccaaagct ggcacggagg aaacaatcct gtactcagac 780
atagacctga agaagctggc tgaaattcgg cagcaaatcc ccattttaaa acagaaacga 840

gcagacctct atacagtgga atcaaagaag ccttgatatc tgtttcaaaa atgtcaccaa 900
caggatgatg ctctgtcaga tgatcaactc tactacatct ctttttttg gagggagggg 960
ggaacagggc catttcatgt taattctatc aatgatctgt gccacaaggt cccctatttt 1020
aattaaaagt ttcatcttta attaaaatgt gcttggtaac aatgttctag ctcttaacta 1080
gtctgatggt tcctaggcat ttcagtccca agatcctttt gaacaattaa aaactgaagc 1140
ctctaagcat tgtttccatg tgtggtgggc tggtcccatc tgtctgagaa aatgtacatt 1200
taccagaaca ctaattttca tggtgctaat atcccatcaa catgacactt ttaaaacttt 1260
ttattaaaaa ttgttttcat acaataaaaa aa 1292

<210> 10
<211> 1214
<212> DNA
<213> X. laevis

<400> 10
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cttcgcggga caggacaggg tcttaggctc tgccttgtgt ccacacgccc ttgtgcagac 120
tgctatagac tgtgacttta accctgtgtc cggatatagg ggttagaagc ctgagtgcaa 180
tggctggtgc ccacaagccc ctgattgccg tgtgccagat gacttcaacc tctgataagg 240
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ggctctggct ttccctgggg ggatttcatg agaaaggacc caactgggac acggaccaac 480
gcatttccaa ttctcacgtg gttgtggaca acacagggca catagtatcg gtgtaccgca 540
aggctcacct gtttgacgta gacttgcaga atggagtgtc actcagagag agcagttcca 600
ccctccccgg agcagagctt attcgcccca tcacttctcc agcaggaaag attggcctgg 660
gggtgtgtta cgacctccgc ttcccagaat tctccttggc tctggcccaa caaggagcag 720
aacttctcac ttacccttct gccttcaccc tcactactgg tctggcacat tgggaggtgt 780
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tggagcgtgt gaggcgggac atgccggtgt ggaggcaccg caggactgat ctgtatggga 1020
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agnctttccc ctgattgctg aaattcctca atctgtgact gtgaatgaca atgaacgtga 1140
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gctgaacctt aaaa 1214

<210> 11
<211> 346
<212> PRT
<213> A. thaliana

<400> 11
Met Ser Ser Thr Lys Asp Met Ser Thr Val Gln Asn Ala Thr Pro Phe
 1   5   10   15
Asn Gly Val Ala Pro Ser Thr Thr Val Arg Val Thr Ile Val Gln Ser
            20   25   30
Ser Thr Val Tyr Asn Asp Thr Pro Ala Thr Ile Asp Lys Ala Glu Lys
        35   40   45
Tyr Ile Val Glu Ala Ala Ser Lys Gly Ala Glu Leu Val Leu Phe Pro
    50   55   60
Glu Gly Phe Ile Gly Gly Tyr Pro Arg Gly Phe Arg Phe Gly Leu Ala
65   70   75   80
Val Gly Val His Asn Glu Glu Gly Arg Asp Glu Phe Arg Lys Tyr His
                85   90   95

Ala Ser Ala Ile His Val Pro Gly Pro Glu Val Ala Arg Leu Ala Asp
100 105 110
Val Ala Arg Lys Asn His Val Tyr Leu Val Met Gly Ala Ile Glu Lys
115 120 125
Glu Gly Tyr Thr Leu Tyr Cys Thr Val Leu Phe Phe Ser Pro Gln Gly
130 135 140
Gln Phe Leu Gly Lys His Arg Lys Leu Met Pro Thr Ser Leu Glu Arg
145 150 155 160
Cys Ile Trp Gly Gln Gly Asp Gly Ser Thr Ile Pro Val Tyr Asp Thr
165 170 175
Pro Ile Gly Lys Leu Gly Ala Ala Ile Cys Trp Glu Asn Arg Met Pro
180 185 190
Leu Tyr Arg Thr Ala Leu Tyr Ala Lys Gly Ile Glu Leu Tyr Cys Ala
195 200 205
Pro Thr Ala Asp Gly Ser Lys Glu Trp Gln Ser Ser Met Leu His Ile
210 215 220
Ala Ile Glu Gly Gly Cys Phe Val Leu Ser Ala Cys Gln Phe Cys Gln
225 230 235 240
Arg Lys His Phe Pro Asp His Pro Asp Tyr Leu Phe Thr Asp Trp Tyr
245 250 255
Asp Asp Lys Glu His Asp Ser Ile Val Ser Gln Gly Gly Ser Val Ile
260 265 270
Ile Ser Pro Leu Gly Gln Val Leu Ala Gly Pro Asn Phe Glu Ser Glu
275 280 285
Gly Leu Val Thr Ala Asp Ile Asp Leu Gly Asp Ile Ala Arg Ala Lys
290 295 300
Leu Tyr Phe Asp Ser Val Gly His Tyr Ser Arg Pro Asp Val Leu His
305 310 315 320
Leu Thr Val Asn Glu His Pro Arg Lys Ser Val Thr Phe Val Thr Lys
325 330 335
Val Glu Lys Ala Glu Asp Asp Ser Asn Lys
340 345